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AMENDMENTS TO THE CLAIMS:

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

1, 2, 6-9, 43, 47 and 52-58. (Cancelled)

- 59. (New) A method for accelerating flowering in a plant, comprising modifying in said plant an endogenous level of at least one compound selected from the group consisting of 12-hydroxyjasmonic acid, sulfate ester of 12-hydroxyjasmonic acid, 11-hydroxyjasmonic acid, sulfate ester of 11-hydroxyjasmonic acid, and mixtures thereof, wherein the endogenous level of the at least one compound is altered by expression of a sulfotransferase encoded by a gene of SEQ ID NO: 1 or a functional homologue having at least 80% similarity to SEQ ID NO:1.
- 60. (New) The method of claim 59, wherein the sulfotransferase has an amino acid sequence of SEQ ID NO: 3 or a functional homologue having at least 80% similarity to SEQ ID NO:3.
- 61. (New) The method of claim 59, wherein said plant is transgenic.
- 62. (New) A method for producing a transgenic plant which flowers early, said method comprising the steps of:
- a) introducing into a cell of a suitable plant an exogenous nucleic acid molecule comprising a sequence of nucleotides antisense to a nucleic acid sequence coding for an amino acid sequence of SEQ ID NO:3 or a functional homologue having at least 80% similarity to SEQ ID NO:3, encoding a plant hydroxyjasmonic acid sulfotransferase:
 - b) regenerating a transgenic plant from the cell; and

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- c) growing said transgenic plant for a time and under conditions sufficient to inhibit expression of the hydroxyjasmonic acid sulfotransferase.
- 63. (New) The method of claim 62, wherein the hydroxyjasmonic acid sulfotransferase is a 11- or a 12- hydroxyjasmonic acid sulfotransferase.
- 64. (New) The method of claim 62, further comprising the step of applying to a plant at least one flowering inducing compounds selected from the group consisting of 12- hydroxyjasmonic acid and 11- hydroxyjasmonic acid.
- 65. (New) The method of claim 62, further comprising the step of applying to said plant at least one inhibitor of a sulfotransferase having an amino acid sequence with at least 80% similarity with SEQ ID NO:3.
- 66. (New) The method of claim 62, further comprising the step of increasing in said plant the endogenous level of an hydroxylase which hydroxylates jasmonic acid or methyljasmonic acid or both.
- 67. (New) The method of claim 62, further comprising the step of inhibiting in said plant the expression of at least one gene selected from the group consisting of SEQ ID NO:1 or a functional homologue having at least 80% similarity to SEQ ID NO::1.
- 68. (New) The method of claim 67, wherein said exogenous sequence is expressed under the control of a constitutive or an inducible promoter.
- 69. (New) A plant genetically modified to flower early wherein the plant is obtained by the method of claim 62.
- 70. (New) The plant of claim 69, which is cauliflower.

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- 71. (New) The plant of claim 69, which is broccoli.
- 72. (New) The plant of claim 69, which is a horticultural plant.
- 73. (New) A cut flower from the genetically modified plant of claim 69.